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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Arabidopsis thaliana ring
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         Submitted (25-AUG-2001) Institute of Plant Biology, Research Center, Temesvari krt. 62, Szeged H-6701, H Location/Qualifiers
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyt
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

[ (bases 1 to 666)
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RIKEN Genomic Sciences Center (GSC) members carried out the collection and clustering of RAFL cDNAs (RAFL CDNA: "RIKEN Arabidopsis Full-Length cDNA'): Seki,M., Narusaka,M., Ishida,J., Satou,M., Kamiya,A., Sakurai,T., Carninci,P., Kawai,J., Hayashizaki,Y. and Shinozaki,K.
                                                                                                        e-mail for correspondence: arab@sequence.stanford.edu
                                                                                                                                        Submitted (02-JAN-2002) DNA Sequencing and Technology Center, Stanford University, 855 California Avenue, Palo Alto, CA 94304,
                                                                                                                                                                                                                                                                                                                                                                                                                 FLI_CDNA
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VCPLDNSEWEFOKYGH"
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The Salk, Stanford, PGEC (SSP) Consortium members carried out the sequencing and annotation of the RAFL cDNAs: Nguyen,M., Southwick,A., Karlin.Neumann,G., Lam,B., Miranda,M., Palm,C.J.,

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Direct Submission
Submitted (04-JUL-1996) Data collected by MIPS on behalf of the
European yeast chromosome XV sequencing project. MIPS at the
                                                                                           Aldea, M., Ca
Unpublished
                                                                                                      Arino, J., Casamayor, A., Gamo, F.J., Aldea, M., Casas, C. and Herrero, E.
                                                                                                                                        Saccharomyces cerevisiae
Eukaryota; Fungli Ascomycota; Saccharomycotina; Saccharomycetas;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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137 c 149 g 19 t
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/chromosome="5"
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                                                                                          Arabidopsis thaliana Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 357)
                           Submitted (25-AUG-2001) Institute o
Research Center, Temesvari krt. 62,
Location/Qualifiers
1. .357
                                                                                Direct Submission
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Martinsried, FRG: E-mail: Mewes@mips.embnet.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           419
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/organism="Arabidopsis
/db_xref="taxon:3702"
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t 209 c 249 g
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/translation="MSNEVDRMDVDEDESQNIAQSSNQSAPVETKKKRPEIKKWTAVAFWSNDIAVDNCAICRNHIMEPCIECQPKAMTDTDNECVAAWGVCNHAFHLHCINKWIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /db_xref="SPTREMBL:008272"
/translation="MARGWHSWHGSIIWFLQIAQLSTAISHDQNATAVHFLISNLFFL
VSTGALWFELCAIFCDSSSTSILSTSILMLNYFSLKFSATGERHLVATNIAHIEAHT
SIVGFMISLFTPLKETIITKYEMLVRT"
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/note="ORF YOL134c"
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/db_xref="GI:1420024"
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/db_xref="taxon:4932"
/chromosome="XV"
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/db_xref="GI:1420023"
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                                                                                                                                                                                                                                                                                                                                                                      The Salk, Stanford, PGEC (SSP) Consortium members carried out th sequencing and annotation of the RAFL CDNAs: Nguyen, M. Southwick, A., Karlin-Neumann, G., Lam, B., Miranda, M., Palm, C.J., Bowser, L., Jones, T., Banh, J., Chen, H., Cheuk, R., Chung, M.K., Kim, C., Lin, J., Liu, S.X., Pham, P.K., Sakano, H., Shinn, P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 666)

Nguyen, M., Karlin-Neumann, G., Southwick, A., Lam, B., Miranda, M., Palm, C.J., Bowser, L., Jones, T., Banh, J., Carninci, P., Chen, H., Cheuk, R., Chung, M.K., Hayashizaki, Y., Ishida, J., Kamiya, A., Kawai, J., Kim, C., Lin, J., Liu, S.X., Narusaka, M., Pham, P.K., Sakano, H., Sakurai, T., Satou, M., Seki, M., Shinn, P., Yamada, K., Shinozaki, K., Ecker, J., Theologis, A. and Davis, R.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RIKEN Genomic Sciences Center (GSC) members carried out the collection and clustering of RAFL cDNAs (RAFL cDNA: 'RIKEN Arabidopsis Full-Length cDNA'): Seki,M., Narusaka,M., Ishida,J., Sakurai,T., Carninci,P., Kawai,J., us., Sakurai,T., Carninci,P., Kawai,J.,
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                                                                                                                                                                                                                                                 Nguyen,M, (SSP/Stanford) and Seki,M. (RIKEN GSC) contrib
equally to this work. Shinozaki,K. (RIKEN GSC) and Davi
(SSP/Stanford) contributed equally to this work as PIs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hayashizaki, Y. and Shinozaki, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e-mail for correspondence: arab@sequence.stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (02-JAN-2002) DNA Sequencing and Technology Center, Stanford University, 855 California Avenue, Palo Alto, CA 94304,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arabidopsis thaliana.
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/gene="At5g20570"
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/product="ring box-1-like protein"
/protein_id="AAALI3435.1"
/protein_id="AAALI3435.1"
/db_xref="GI:16186265"
/translation="MATIDSDYMIPAGEASSSVAASSSNKKAKRFEIKKWSAVALWAWIVUNCALICIECQANQASATSEECTVAMGVCNHAFHFHCISRWLKTRQVCPLDNSEWEFQKYGH"
                                                         /note-"This clone is in ecotype: Columbia"
                                                                                                                                           /chromosome="5"
                                                                                                                                                             /organism="Arabidopsis thaliana"
/db_xref="taxon:3702"
                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                /clone="RAFL09-78-L12"
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712 AAAAAGGAACATGCCATAATCCTTCTACCTCGGTATGATTTAAATGTTTACGG
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2 /h--
                                                               l Similarity
28; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     European yeast chromosome XV sequencing project. MIPS at Max-Planck-Institut fuer Blochemie, Am Klopferspitz 18a Martinsried, FRG; E-mail: Mewes@mips.embnet.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S.cerevisiae (
274876 Y13140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (04-JUL-1996) Data collected by MIPS on behalf of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Fungi; Ascomycota; Saccharon Saccharomycetales; Saccharomycetaceae; 1 (bases 1 to 1255)
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